

(#11)

50C

## RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/523,893A  
Source: PCP/10  
Date Processed by STIC: 1/9/06

# *ENTERED*

# CRF Errors Edited by the STIC Systems Branch

Serial Number: 10/523,893A

CRF Edit Date: 1/11/06  
Edited by: AS

— Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

— Corrected the SEQ ID NO. Sequence numbers edited were:

— Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

— Deleted: \_\_\_\_\_ invalid beginning/end-of-file text ; \_\_\_\_\_ page numbers

— Inserted mandatory headings/numeric identifiers, specifically:

— Moved responses to same line as heading/numeric identifier, specifically:

— Other:



RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/523,893A

DATE: 01/11/2006  
TIME: 12:16:19

Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\01112006\J523893A.raw

3 <110> APPLICANT: Lowery, David E.  
4 Smith, Valdin G.  
5 Kubiak, Teresa M.  
6 Larsen, Martha J.  
8 <120> TITLE OF INVENTION: Drosophila G Protein Coupled Receptors, Nucleic Acids, and  
9 Methods Related to the Same  
11 <130> FILE REFERENCE: PHRM0002-105  
13 <140> CURRENT APPLICATION NUMBER: US 10/523,893A  
14 <141> CURRENT FILING DATE: 2005-02-04  
16 <150> PRIOR APPLICATION NUMBER: US 10/283,423  
17 <151> PRIOR FILING DATE: 2002-10-30  
19 <150> PRIOR APPLICATION NUMBER: US 09/693,746  
20 <151> PRIOR FILING DATE: 2000-10-20  
22 <150> PRIOR APPLICATION NUMBER: US 09/425,676  
23 <151> PRIOR FILING DATE: 1999-10-22  
25 <160> NUMBER OF SEQ ID NOS: 232  
27 <170> SOFTWARE: PatentIn version 3.3  
29 <210> SEQ ID NO: 1  
30 <211> LENGTH: 1803  
31 <212> TYPE: DNA  
32 <213> ORGANISM: D. melanogaster  
34 <400> SEQUENCE: 1  
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37 cagctccat tggcggcac aaccaactgg agcctaacgt cgccggaaac tactagcgct 120  
39 atcttggccgg atgtggctgc atcggatgag gataggagcg gcgggatcat tcacaaccag 180  
41 ttcgtgcaaa tcttcttcta cgtcctgtac gcccacggct ttgtcctggg tgcgttcgg 240  
43 aatgtccctgg tttgctacgt agttctgagg aatcgggcca tgcagactgt gaccaatata 300  
45 ttcatcacga atctggccct gtcggacata ttgctctgcg tcctggcggt gccattttact 360  
47 ccgcttaca cgttcatggg tcgctgggcc ttccggcggg gtctgtgcca tctgggtgtcc 420  
49 ttggcccgagg gatgcagcat ctacatatcc acgctgaccc tcacccatcgat tgccatcgat 480  
51 cggtaacctcg ttatcatata cccctccat cccgcgcata agcttccac ctgcacatggg 540  
53 atcatagtga gcatctgggt gatagccctg ctggccaccc ttccctacgg catgtacatg 600  
55 aagatgacca acgagctggta gaacggaaacg cagacaggca acgagaccct ggtggaggcc 660  
57 actctaattgc taaacggaaat ctttggcc caggatcgat gattcatcgat ggcggccggac 720  
59 tctacccatgg ccacccaggc ctatatcgat gtgatgaccc cggatcaac gggaccggag 780  
61 atgcctatg tgggggtgtta ctgcggaggag aactggccat cggagcgtta cccggaaagggt 840  
63 ttcggtgcca tcacaaaccac tctgcagttt gtgtccctt tcttcatcat ctcgatttc 900  
65 tacgtgtgaa tatcggtgaa gctaaaccag cggggccaggc ccaagccggg atcgaaatcc 960  
67 tcgagacggg aggaggcgga tcggatcgca aagaagcgca ccaaccgcgt gtcacatcgcc 1020  
69 atgggtggccgg tattcgact cagctggctg cccatcaatg tggtaacat attcgatgac 1080  
71 ttcgatgaca agtccaacga gtggcgcttc tacatcctat tcttctttgt ggcccactct 1140  
73 attgcccattga gctccacccctg ctacaatccc ttcctgtacg cctggctgaa cgagaacttc 1200  
75 cgcaaggagt tcaagcactg gctgcctgc ttaatccct cgaacaacaa catcatcaac 1260

P.6

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/523,893A

DATE: 01/11/2006  
TIME: 12:16:19

Input Set : A:\PTO.AMC.txt  
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77 atcaccaggg gctataatcg gagtgatcg aacacctgtg gtccgcgact gcatcatggc 1320  
 79 aaggggatg gtggcatggg cggtggcagt ctggacgccc acgaccagga cgagaacggc 1380  
 81 atcacccagg agacctgtct gccccaggag aagctgctga ttatccccag ggagccgact 1440  
 83 tacggcaatg gcacgggtgc cgtgtcgcata ctccttagcg ggcgcggcat taacgcgc 1500  
 85 ctggtgacacg gtggcggacca tcagatgcac cagctgcagc cgtcacacca tcaacaggtg 1560  
 87 gagctgacga ggcaaatccg cccggcggaca gacgagacgg acggggattt cctggactcc 1620  
 89 ggcgacgagc agaccgtgga ggtgcgccttc agcgagacgc cggtcgtagc cacggataat 1680  
 91 accaccggga tcagcattct ggagacgagt acgagtcaact gccaggactc ggtatgtatg 1740  
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 95 tga 1803  
 98 <210> SEQ ID NO: 2  
 99 <211> LENGTH: 600  
 100 <212> TYPE: PRT  
 101 <213> ORGANISM: D melanogaster  
 103 <400> SEQUENCE: 2  
 105 Met Ala Asn Leu Ser Trp Leu Ser Thr Ile Thr Thr Thr Ser Ser Ser  
 106 1 5 10 15  
 109 Ile Ser Thr Ser Gln Leu Pro Leu Val Ser Thr Thr Asn Trp Ser Leu  
 110 20 25 30  
 113 Thr Ser Pro Gly Thr Thr Ser Ala Ile Leu Ala Asp Val Ala Ala Ser  
 114 35 40 45  
 117 Asp Glu Asp Arg Ser Gly Gly Ile Ile His Asn Gln Phe Val Gln Ile  
 118 50 55 60  
 121 Phe Phe Tyr Val Leu Tyr Ala Thr Val Phe Val Leu Gly Val Phe Gly  
 122 65 70 75 80  
 125 Asn Val Leu Val Cys Tyr Val Val Leu Arg Asn Arg Ala Met Gln Thr  
 126 85 90 95  
 129 Val Thr Asn Ile Phe Ile Thr Asn Leu Ala Leu Ser Asp Ile Leu Leu  
 130 100 105 110  
 133 Cys Val Leu Ala Val Pro Phe Thr Pro Leu Tyr Thr Phe Met Gly Arg  
 134 115 120 125  
 137 Trp Ala Phe Gly Arg Ser Leu Cys His Leu Val Ser Phe Ala Gln Gly  
 138 130 135 140  
 141 Cys Ser Ile Tyr Ile Ser Thr Leu Thr Leu Thr Ser Ile Ala Ile Asp  
 142 145 150 155 160  
 145 Arg Tyr Phe Val Ile Ile Tyr Pro Phe His Pro Arg Met Lys Leu Ser  
 146 165 170 175  
 149 Thr Cys Ile Gly Ile Ile Val Ser Ile Trp Val Ile Ala Leu Leu Ala  
 150 180 185 190  
 153 Thr Val Pro Tyr Gly Met Tyr Met Lys Met Thr Asn Glu Leu Val Asn  
 154 195 200 205  
 157 Gly Thr Gln Thr Gly Asn Glu Thr Leu Val Glu Ala Thr Leu Met Leu  
 158 210 215 220  
 161 Asn Gly Ser Phe Val Ala Gln Gly Ser Gly Phe Ile Glu Ala Pro Asp  
 162 225 230 235 240  
 165 Ser Thr Ser Ala Thr Gln Ala Tyr Met Gln Val Met Thr Ala Gly Ser  
 166 245 250 255  
 169 Thr Gly Pro Glu Met Pro Tyr Val Arg Val Tyr Cys Glu Glu Asn Trp  
 170 260 265 270

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/523,893A

DATE: 01/11/2006  
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Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\01112006\J523893A.raw

173 Pro Ser Glu Gln Tyr Arg Lys Val Phe Gly Ala Ile Thr Thr Thr Leu  
174 275 280 285  
177 Gln Phe Val Leu Pro Phe Phe Ile Ile Ser Ile Cys Tyr Val Trp Ile  
178 290 295 300  
181 Ser Val Lys Leu Asn Gln Arg Ala Arg Ala Lys Pro Gly Ser Lys Ser  
182 305 310 315 320  
185 Ser Arg Arg Glu Glu Ala Asp Arg Asp Arg Lys Lys Arg Thr Asn Arg  
186 325 330 335  
189 Met Leu Ile Ala Met Val Ala Val Phe Gly Leu Ser Trp Leu Pro Ile  
190 340 345 350  
193 Asn Val Val Asn Ile Phe Asp Asp Phe Asp Asp Lys Ser Asn Glu Trp  
194 355 360 365  
197 Arg Phe Tyr Ile Leu Phe Phe Val Ala His Ser Ile Ala Met Ser  
198 370 375 380  
201 Ser Thr Cys Tyr Asn Pro Phe Leu Tyr Ala Trp Leu Asn Glu Asn Phe  
202 385 390 395 400  
205 Arg Lys Glu Phe Lys His Val Leu Pro Cys Phe Asn Pro Ser Asn Asn  
206 405 410 415  
209 Asn Ile Ile Asn Ile Thr Arg Gly Tyr Asn Arg Ser Asp Arg Asn Thr  
210 420 425 430  
213 Cys Gly Pro Arg Leu His His Gly Lys Gly Asp Gly Gly Met Gly Gly  
214 435 440 445  
217 Gly Ser Leu Asp Ala Asp Asp Gln Asp Glu Asn Gly Ile Thr Gln Glu  
218 450 455 460  
221 Thr Cys Leu Pro Lys Glu Lys Leu Leu Ile Ile Pro Arg Glu Pro Thr  
222 465 470 475 480  
225 Tyr Gly Asn Gly Thr Gly Ala Val Ser Pro Ile Leu Ser Gly Arg Gly  
226 485 490 495  
229 Ile Asn Ala Ala Leu Val His Gly Gly Asp His Gln Met His Gln Leu  
230 500 505 510  
233 Gln Pro Ser His His Gln Gln Val Glu Leu Thr Arg Arg Ile Arg Arg  
234 515 520 525  
237 Arg Thr Asp Glu Thr Asp Gly Asp Tyr Leu Asp Ser Gly Asp Glu Gln  
238 530 535 540  
241 Thr Val Glu Val Arg Phe Ser Glu Thr Pro Phe Val Ser Thr Asp Asn  
242 545 550 555 560  
245 Thr Thr Gly Ile Ser Ile Leu Glu Thr Ser Thr Ser His Cys Gln Asp  
246 565 570 575  
249 Ser Asp Val Met Val Glu Leu Gly Glu Ala Ile Gly Ala Gly Gly Gly  
250 580 585 590  
253 Ala Glu Leu Gly Arg Arg Ile Asn  
254 595 600  
257 <210> SEQ ID NO: 3  
258 <211> LENGTH: 1445  
259 <212> TYPE: DNA  
260 <213> ORGANISM: D. melanogaster  
262 <400> SEQUENCE: 3  
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265 agcagcaaca gcgtgcgcta tctggacgac cggcatccgc tggactacct tgacctggc 120

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Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\01112006\J523893A.raw

267	acgggtgcacg	ccctcaacac	cactgccatc	aacacacctgg	atctgaatga	gactgggagc	180
269	aggccgctgg	accccggtgc	tatcgatagg	ttcctgagca	acagggcggt	ggacagcccc	240
271	tggtaccaca	tgtctcatca	catgtacggc	gtgctaattcg	tcttcggcgc	ccttaggcaac	300
273	accctggttt	ttatagccgt	catccggaag	cccatcatgc	gcactgctcg	caatctttc	360
275	atcctcaacc	tggccatata	ggacctactt	ttatgcctag	tcaccatgcc	gctgaccctt	420
277	atggagatcc	tgtccaaagta	ctggccctac	ggctcctgt	ccatcctgtg	caaaaacgatt	480
279	gccatgctgc	aggcactttg	tatttcgtg	tcgacaat	ccataacggc	cattgccttc	540
281	gacagatatac	aggtgtatcg	gtaccccacg	cgggacagcc	tgcagttcg	gggcgcgtg	600
283	acgatctgg	cggggatctg	ggcactggca	ctgctgttgg	cctcggcgt	gttctgttac	660
285	aaggagctga	tcaacacaga	cacgcccggca	ctcctgcagc	agatggcct	gcaggacacg	720
287	atccctgtact	gcatttgagga	ctggccaagt	cgcaacgggc	gcttctacta	ctcgatcttc	780
289	tcgctgtcg	tacaataac	gtgtccatc	ctgatcgct	cggtggcata	cttcgggatc	840
291	tacaacaagc	tgaagagccg	catcaccgtg	gtggctgtc	aggcgtcctc	cgctcagcgg	900
293	aagggtggagc	ggggggcggcg	gatgaagcgc	accaactgcc	tactgatcg	catcgccatc	960
295	atctttggcg	tttcttggct	gccgctgaac	tttttcaacc	tgtacgcgg	catggagcgc	1020
297	tcgcccgtca	ctcagagcat	gctagccgc	tacgccatct	gccacatgt	cggcgtgagc	1080
299	tccgcctgt	ccaacccgtt	gctctacggc	tggctcaacg	acaacttccg	taaagaattt	1140
301	caagaactgc	tctgcccgtt	ctcagacact	aatgttgctc	ttaacggtca	cacgacagggc	1200
303	tgcaacgtcc	aggcggcggc	gchgcaagcgt	cgcaagttgg	gcccggaaact	ctccaaaggc	1260
305	gaactcaagc	tgtctggggcc	aggcggcggc	cagagcggta	ccgcggcgggg	ggaaggcgg	1320
307	ctggcggcca	ccgacttcat	gaccggccac	cacgaggggc	gactgcgcag	cgccataacc	1380
309	gagtcgggtgg	ccctcagcgg	ccacaacccc	gtgcctcgg	aggtcaccaa	gctgatgccg	1440
311	cggtta						1445

314 <210> SEQ ID NO: 4

315 <211> LENGTH: 357

316 <212> TYPE: PRT

317 <213> ORGANISM: D. melanogaster

319 <400> SEQUENCE: 4

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322	1					5				10					15	
325	Asn	Glu	Glu	Asn	Ile	Thr	Ser	Phe	Phe	Thr	Asp	Glu	Glu	Trp	Leu	Ala
326						20				25				30		
329	Ile	Asn	Gly	Thr	Leu	Pro	Trp	Ile	Val	Gly	Phe	Phe	Phe	Gly	Val	Ile
330						35				40				45		
333	Ala	Ile	Thr	Gly	Phe	Phe	Gly	Asn	Leu	Leu	Val	Ile	Leu	Val	Val	Val
334						50				55				60		
337	Phe	Asn	Asn	Asn	Met	Arg	Ser	Thr	Thr	Asn	Leu	Met	Ile	Val	Asn	Leu
338						65				70				75		80
341	Ala	Ala	Ala	Asp	Leu	Met	Phe	Val	Ile	Leu	Cys	Ile	Pro	Phe	Thr	Ala
342						85				90				95		
345	Thr	Asp	Tyr	Met	Val	Tyr	Tyr	Trp	Pro	Tyr	Gly	Arg	Phe	Trp	Cys	Arg
346						100				105				110		
349	Ser	Val	Gln	Tyr	Leu	Ile	Val	Val	Thr	Ala	Phe	Ala	Ser	Ile	Tyr	Thr
350						115				120				125		
353	Leu	Val	Leu	Met	Ser	Ile	Asp	Arg	Phe	Leu	Ala	Val	Val	His	Pro	Ile
354						130				135				140		
357	Arg	Ser	Arg	Met	Met	Arg	Thr	Glu	Asn	Ile	Thr	Leu	Ile	Ala	Ile	Val
358	145					150				155				160		
361	Thr	Leu	Trp	Ile	Val	Val	Leu	Val	Val	Ser	Val	Pro	Val	Ala	Phe	Thr

RAW SEQUENCE LISTING  
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Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\01112006\J523893A.raw

362	165	170	175	
365	His Asp Val Val Val Asp Tyr Asp Ala Lys Lys Asn Ile Thr Tyr Gly			
366	180	185	190	
369	Met Cys Thr Phe Thr Thr Asn Asp Phe Leu Gly Pro Arg Thr Tyr Gln			
370	195	200	205	
373	Val Thr Phe Phe Ile Ser Ser Tyr Leu Leu Pro Leu Met Ile Ile Ser			
374	210	215	220	
377	Gly Leu Tyr Met Arg Met Ile Met Arg Leu Trp Arg Gln Gly Thr Gly			
378	225	230	235	240
381	Val Arg Met Ser Lys Glu Ser Gln Arg Gly Arg Lys Arg Val Thr Arg			
382	245	250	255	
385	Leu Val Val Val Val Val Ile Ala Phe Ala Ser Leu Trp Leu Pro Val			
386	260	265	270	
389	Gln Leu Ile Leu Leu Lys Ser Leu Asp Val Ile Glu Thr Asn Thr			
390	275	280	285	
393	Leu Thr Lys Leu Val Ile Gln Val Thr Ala Gln Thr Leu Ala Tyr Ser			
394	290	295	300	
397	Ser Ser Cys Ile Asn Pro Leu Leu Tyr Ala Phe Leu Ser Glu Asn Phe			
398	305	310	315	320
401	Arg Lys Ala Phe Tyr Lys Ala Val Asn Cys Ser Ser Arg Tyr Gln Asn			
402	325	330	335	
405	Tyr Thr Ser Asp Leu Pro Pro Pro Arg Lys Thr Ser Cys Ala Arg Thr			
406	340	345	350	
409	Ser Thr Thr Gly Leu			
410	355			
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414	<211> LENGTH: 1376			
415	<212> TYPE: DNA			
416	<213> ORGANISM: D. melanogaster			
418	<400> SEQUENCE: 5			
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421	agcagcaaca gcgtgcgcta tctggacgac cggcatccgc tggactacct tgacctggc	120		
423	acggcgcacg ccctcaacac cactgccatc aacacctcg atctgaatga gactggagc	180		
425	aggccgctgg accccggtct tatcgatagg ttccctgagca acaggccggt ggacagcccc	240		
427	tggtaccaca tgctcatcg catgtacggc gtgctaatcg tcttcggcgc cctaggcaac	300		
429	accctggttg ttatagccgt catccggaag cccatcatgc gcactgctcg caatctttc	360		
431	atcctaacc tggccatatac ggacctactt ttatgcctag tcaccatgcc gctgaccc	420		
433	atggagatcc tgcctcaagta ctggccctac ggctcctgtc ccatctgtg caaaacgatt	480		
435	gccatgctgc aggcactttg tatttcgtg tcgacaatat ccataacggc cattgccttc	540		
437	gacagatatac aggtgatcgt gtacccacg cgggacagcc tgcagttcg gggcgcggtg	600		
439	acgatcctgg cggggatctg ggcactggca ctgctgctgg cctccgcgt gttcgctac	660		
441	aaggagctga tcaacacaga cacgcggca ctcctgcagc agatcgccgc gcaggacacg	720		
443	atcccgtaact gcattgagga ctggccaagt cgcacacggc gcttctacta ctcgatctc	780		
445	tcgctgtgcg tacaataacct ggtgccatc ctgatcgct cgggtggcata cttcgggatc	840		
447	tacaacaagg tgaagagccg catcaccgtg gtggctgtgc aggctgcctc cgctcagcgg	900		
449	aaggtggagc gggggcggcg gatgaagcgc accaactgcc tactgatcag catgcacatc	960		
451	atcttggcg ttcttggct gccgctgaac ttttcaacc tgtacgcggc catggagcgc	1020		
453	tcgcgggtca ctcagagcat gctagtccgc tacgccatct gccacatgat cggcatgagc	1080		
455	tccgcctgtc ccaaccgtt gctctacggc tggctcaacg acaacttccg ctgcaacgtc	1140		

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/523,893A

DATE: 01/11/2006  
TIME: 12:16:20

Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\01112006\J523893A.raw

Base Note:

z of n and/or Xaa have been detected in the Sequence Listing. Please review the  
Sequence Listing to ensure that a corresponding explanation is presented in the <220>  
<223> fields of each sequence which presents at least one n or Xaa.

z#:50; Xaa Pos. 3  
z#:88; Xaa Pos. 1  
z#:120; Xaa Pos. 1  
z#:138; Xaa Pos. 1  
z#:151; Xaa Pos. 1  
z#:177; Xaa Pos. 1  
z#:182; Xaa Pos. 1  
z#:184; Xaa Pos. 1  
z#:185; Xaa Pos. 1  
z#:232; Xaa Pos. 1

**VERIFICATION SUMMARY** DATE: 01/11/2006  
PATENT APPLICATION: US/10/523,893A TIME: 12:16:20

Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\01112006\J523893A.raw

3675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:0  
3213 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:88 after pos.:0  
3667 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:120 after pos.:0  
3923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138 after pos.:0  
4111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:151 after pos.:0  
4479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177 after pos.:0  
4555 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:182 after pos.:0  
4589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:184 after pos.:0  
4609 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:185 after pos.:0  
5189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:232 after pos.:0

**Raw Sequence Listing before editing,  
for reference only**



PCT

RAW SEQUENCE LISTING DATE: 01/09/2006  
 PATENT APPLICATION: US/10/523,893A TIME: 12:05:37

Input Set : A:\PHRM0002-105.ST25.txt  
 Output Set: N:\CRF4\01092006\J523893A.raw

3 <110> APPLICANT: Lowery, David E.  
 4 Smith, Valdin G.  
 5 Kubiak, Teresa M.  
 6 Larsen, Martha J.  
 8 <120> TITLE OF INVENTION: Drosophila G Protein Coupled Receptors, Nucleic Acids, and  
 9 Methods Related to the Same  
 11 <130> FILE REFERENCE: PHRM0002-105  
 13 <140> CURRENT APPLICATION NUMBER: US 10/523,893A  
 14 <141> CURRENT FILING DATE: 2005-02-04  
 16 <150> PRIOR APPLICATION NUMBER: US 10/283,423  
 17 <151> PRIOR FILING DATE: 2002-10-30  
 19 <150> PRIOR APPLICATION NUMBER: US 09/693,746  
 20 <151> PRIOR FILING DATE: 2000-10-20  
 22 <150> PRIOR APPLICATION NUMBER: US 09/425,676  
 23 <151> PRIOR FILING DATE: 1999-10-22  
 25 <160> NUMBER OF SEQ ID NOS: 232  
 27 <170> SOFTWARE: PatentIn version 3.3

## ERRORED SEQUENCES

5173 <210> SEQ ID NO: 232  
 5174 <211> LENGTH: 14  
 5175 <212> TYPE: PRT  
 5176 <213> ORGANISM: Artificial Sequence  
 5178 <220> FEATURE:  
 5179 <223> OTHER INFORMATION: Novel Sequence  
 5182 <220> FEATURE:  
 5183 <221> NAME/KEY: MOD\_RES  
 5184 <222> LOCATION: (1)..(1)  
 5185 <223> OTHER INFORMATION: Xaa is pGlu  
 5187 <400> SEQUENCE: 232  
 W--> 5189 Xaa Val Arg Phe Gln Cys Tyr Phe Asn Pro Ile Ser Cys Phe  
 5190 1 5 10  
 E--> 5193 - 1 -  
 E--> 5195 - 17 -

*Does Not Comply  
 Corrected Diskette Needed*

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/523,893A

DATE: 01/09/2006

TIME: 12:05:38

Input Set : A:\PHRM0002-105.ST25.txt  
Output Set: N:\CRF4\01092006\J523893A.raw

L:2675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:0  
L:3213 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:88 after pos.:0  
L:3667 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:120 after pos.:0  
L:3923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138 after pos.:0  
L:4111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:151 after pos.:0  
L:4479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177 after pos.:0  
L:4555 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:182 after pos.:0  
L:4589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:184 after pos.:0  
L:4609 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:185 after pos.:0  
L:5189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:232 after pos.:0  
L:5193 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:232  
L:5195 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:232